

**BLAST 2 SEQUENCES RESULTS**

Query: SEQ ID NO:1 of Serial No. 60/216,523

Subject: SEQ ID NO:1 of Serial No. 09/899,532

Score = 3288 bits (1710), Expect = 0.0

Identities = 1710/1710 (100%)

Strand = Plus / Plus

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Query: 1      tggccctcgaggccaagaattcggcacgaggaggcggggagccagaggcgccaggaccct 60
              |||||||||||||||||||||||||||||||||||||||||||||||||||
Sbjct: 1      tggccctcgaggccaagaattcggcacgaggaggcggggagccagaggcgccaggaccct 60

Query: 61     cgcggtggcgctccagcaccacagaccgtggcgggcgccctcgcccttagggaagagcaaggga 120
              |||||||||||||||||||||||||||||||||||||||||||||||||||
Sbjct: 61     cgcggtggcgctccagcaccacagaccgtggcgggcgccctcgcccttagggaagagcaaggga 120

Query: 121    agaactttatttgaaccgcgaacattttttggtcactgagatcgagtctcccagtgcttt 180
              |||||||||||||||||||||||||||||||||||||||||||||||||||
Sbjct: 121    agaactttatttgaaccgcgaacattttttggtcactgagatcgagtctcccagtgcttt 180

Query: 181    ggcttcccgcctctttatcggtgggtttgatccctgagctgctctcctttcccgaacctcc 240
              |||||||||||||||||||||||||||||||||||||||||||||||||||
Sbjct: 181    ggcttcccgcctctttatcggtgggtttgatccctgagctgctctcctttcccgaacctcc 240

Query: 241    cggggtgcagcctagagccctcccgcgcggctgactccagagtagaggaagggaggcggc 300
              |||||||||||||||||||||||||||||||||||||||||||||||||||
Sbjct: 241    cggggtgcagcctagagccctcccgcgcggctgactccagagtagaggaagggaggcggc 300

Query: 301    ctccggctggtccccgaagccctcgctgccccgcagatgcggatggccagccagtagcg 360
              |||||||||||||||||||||||||||||||||||||||||||||||||||
Sbjct: 301    ctccggctggtccccgaagccctcgctgccccgcagatgcggatggccagccagtagcg 360

Query: 361    ggcggtggccccgcgtcccgggagcgcacagcaatgcaggcgcttaacattaccccgag 420
              |||||||||||||||||||||||||||||||||||||||||||||||||||
Sbjct: 361    ggcggtggccccgcgtcccgggagcgcacagcaatgcaggcgcttaacattaccccgag 420

Query: 421    cagttctctcggctgctgcgggaccacaacctgacgcgggagcagttcatcgctctgtac 480
              |||||||||||||||||||||||||||||||||||||||||||||||||||
Sbjct: 421    cagttctctcggctgctgcgggaccacaacctgacgcgggagcagttcatcgctctgtac 480
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Query: 481 **cggtcgacgctcgtctacaccccagagctgccgggacgcgccaagctggccctcgtg** 540  
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 Sbjct: 481 **cggtcgacgctcgtctacaccccagagctgccgggacgcgccaagctggccctcgtg** 540

Query: 541 **ctcacggcggtgctcatcttcgccctggcgctctttggcaatgctctggtgttctacgtg** 600  
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 Sbjct: 541 **ctcacggcggtgctcatcttcgccctggcgctctttggcaatgctctggtgttctacgtg** 600

Query: 601 **gtgaccgcagcaaggccatgcgccacgtcaccaacatctttatctgctccttggcgctc** 660  
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 Sbjct: 601 **gtgaccgcagcaaggccatgcgccacgtcaccaacatctttatctgctccttggcgctc** 660

Query: 661 **agtgacctgctcatcaccttcttctgcattcccgtcaccatgctccagaacatttccgac** 720  
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 Sbjct: 661 **agtgacctgctcatcaccttcttctgcattcccgtcaccatgctccagaacatttccgac** 720

Query: 721 **aactggctggggggtgctttcatttgcaagatggtgccatttgtccagtctaccgctgtt** 780  
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 Sbjct: 721 **aactggctggggggtgctttcatttgcaagatggtgccatttgtccagtctaccgctgtt** 780

Query: 781 **gtgacagaaatcctcactatgacctgcattgctgtggaaaggcaccagggacttgtgcat** 840  
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 Sbjct: 781 **gtgacagaaatcctcactatgacctgcattgctgtggaaaggcaccagggacttgtgcat** 840

Query: 841 **ccttttaaaatgaagtggcaatacaccaaccgaagggtttcacaatgctaggtgtggtc** 900  
 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
 Sbjct: 841 **ccttttaaaatgaagtggcaatacaccaaccgaagggtttcacaatgctaggtgtggtc** 900

Query: 901 **tggctggtggcagtcacgttaggatcacccatgtggcacgtgcaacaacttgagatcaaa** 960  
 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
 Sbjct: 901 **tggctggtggcagtcacgttaggatcacccatgtggcacgtgcaacaacttgagatcaaa** 960

Query: 961 **tatgacttcctatatgaaaaggaacacatctgctgcttagaagagtggaccagccctgtg** 1020  
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 Sbjct: 961 **tatgacttcctatatgaaaaggaacacatctgctgcttagaagagtggaccagccctgtg** 1020

Query: 1021 **caccagaagatctacaccaccttcaccttgtcatcctcttcctcctgcctcttatgggtg** 1080  
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 Sbjct: 1021 **caccagaagatctacaccaccttcaccttgtcatcctcttcctcctgcctcttatgggtg** 1080

Query: 1081 **atgcttattctgtacagtaaaattgggttatgaactttggataaagaaaagagtggggat** 1140  
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 Sbjct: 1081 **atgcttattctgtacagtaaaattgggttatgaactttggataaagaaaagagtggggat** 1140

Query: 1141 **ggttcagtgcttcgaactattcatggaaaagaaatgtccaaaatagccaggaagaagaaa** 1200  
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 Sbjct: 1141 **ggttcagtgcttcgaactattcatggaaaagaaatgtccaaaatagccaggaagaagaaa** 1200

Query: 1201 **cgagctgtcattatgatgggtgacagtggtggctctctttgctgtgtgctgggcaccattc** 1260  
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 Sbjct: 1201 **cgagctgtcattatgatgggtgacagtggtggctctctttgctgtgtgctgggcaccattc** 1260

Query: 1261 **catgttggtccatgatgatgattgaatacagtaattttgaaaaggaatatgatgatgtcaca** 1320  
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 Sbjct: 1261 **catgttggtccatgatgatgattgaatacagtaattttgaaaaggaatatgatgatgtcaca** 1320

Query: 1321 **atcaagatgatttttgctatcgtgcaaattattggattttccaactccatctgtaatccc** 1380  
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 Sbjct: 1321 **atcaagatgatttttgctatcgtgcaaattattggattttccaactccatctgtaatccc** 1380

Query: 1381 **attgtctatgcatttatgaatgaaaacttcaaaaaaatgttttgcctgcagtttggtat** 1440  
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 Sbjct: 1381 **attgtctatgcatttatgaatgaaaacttcaaaaaaatgttttgcctgcagtttggtat** 1440

Query: 1441 **tgcatagtaaataaaaaccttctctccagcacaaaggcatggaaattcaggaattacaatg** 1500  
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 Sbjct: 1441 **tgcatagtaaataaaaaccttctctccagcacaaaggcatggaaattcaggaattacaatg** 1500

Query: 1501 **atgcggaagaaagcaaagttttccctcagagagaatccagtggaggaaaccaaaggagaa** 1560  
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 Sbjct: 1501 **atgcggaagaaagcaaagttttccctcagagagaatccagtggaggaaaccaaaggagaa** 1560

Query: 1561 **gcattcagtgatggcaacattgaagtcaaattgtgtgaacagacagaggagaagaaaaag** 1620  
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 Sbjct: 1561 **gcattcagtgatggcaacattgaagtcaaattgtgtgaacagacagaggagaagaaaaag** 1620

Query: 1621 **ctcaaacgacatcttgctctcttttaggtctgaactggctgagaattctccttttagacagt** 1680  
 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
 Sbjct: 1621 **ctcaaacgacatcttgctctcttttaggtctgaactggctgagaattctccttttagacagt** 1680

Query: 1681 **gggcattaattataacaatatcttcataat** 1710  
 ||||||||||||||||||||||||||||  
 Sbjct: 1681 **gggcattaattataacaatatcttcataat** 1710

**BLAST 2 SEQUENCES RESULTS**

Query: SEQ ID NO:2 of Serial No. 60/216,523

Subject: SEQ ID NO:2 of Serial No. 09/899,532

Score = 865 bits (2235), Expect = 0.0  
Identities = 431/431 (100%), Positives = 431/431 (100%)

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Query: 1  MQALNITPEQFSRLLRDHNLTREQFIALYRLRPLVYTPELPGRAKLALVLTGVLIFALAL 60
          MQALNITPEQFSRLLRDHNLTREQFIALYRLRPLVYTPELPGRAKLALVLTGVLIFALAL
Sbjct: 1  MQALNITPEQFSRLLRDHNLTREQFIALYRLRPLVYTPELPGRAKLALVLTGVLIFALAL 60

Query: 61  FGNALVFYVVTRSKAMRTVTNIFICSLALSDLLITFFCIPVTMLQNISDNWLGGAFICKM 120
          FGNALVFYVVTRSKAMRTVTNIFICSLALSDLLITFFCIPVTMLQNISDNWLGGAFICKM
Sbjct: 61  FGNALVFYVVTRSKAMRTVTNIFICSLALSDLLITFFCIPVTMLQNISDNWLGGAFICKM 120

Query: 121 VPFVQSTAVVTEILTMTCIAVERHQGLVHPFKMKWQYTNRRRAFTMLGVVWLVAVIVGSPM 180
          VPFVQSTAVVTEILTMTCIAVERHQGLVHPFKMKWQYTNRRRAFTMLGVVWLVAVIVGSPM
Sbjct: 121 VPFVQSTAVVTEILTMTCIAVERHQGLVHPFKMKWQYTNRRRAFTMLGVVWLVAVIVGSPM 180

Query: 181 WHVQQLEIKYDFLYEKEHICCLEEWTSPPVHQKIYTTFILVILFLLPLMVMLILYSKIGYE 240
          WHVQQLEIKYDFLYEKEHICCLEEWTSPPVHQKIYTTFILVILFLLPLMVMLILYSKIGYE
Sbjct: 181 WHVQQLEIKYDFLYEKEHICCLEEWTSPPVHQKIYTTFILVILFLLPLMVMLILYSKIGYE 240

Query: 241 LWIKKRVGDGSLRRTIHKEMSKIARKKKRAVIMMVTVVALFAVCWAPFHVVHMMIEYSN 300
          LWIKKRVGDGSLRRTIHKEMSKIARKKKRAVIMMVTVVALFAVCWAPFHVVHMMIEYSN
Sbjct: 241 LWIKKRVGDGSLRRTIHKEMSKIARKKKRAVIMMVTVVALFAVCWAPFHVVHMMIEYSN 300

Query: 301 FEKEYDDVTIKMIFAIVQIIGFSNSICNPITYAFMNENFKKNVLSAVCYCIVNKTFSPAQ 360
          FEKEYDDVTIKMIFAIVQIIGFSNSICNPITYAFMNENFKKNVLSAVCYCIVNKTFSPAQ
Sbjct: 301 FEKEYDDVTIKMIFAIVQIIGFSNSICNPITYAFMNENFKKNVLSAVCYCIVNKTFSPAQ 360

Query: 361 RHGNSGITMMRKKAKFSLRENPEETKGEAFSDGNIEVKLCEQTEKKKKLRHLALFRSE 420
          RHGNSGITMMRKKAKFSLRENPEETKGEAFSDGNIEVKLCEQTEKKKKLRHLALFRSE
Sbjct: 361 RHGNSGITMMRKKAKFSLRENPEETKGEAFSDGNIEVKLCEQTEKKKKLRHLALFRSE 420

Query: 421 LAENSPLDSGH 431
          LAENSPLDSGH
Sbjct: 421 LAENSPLDSGH 431

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